

## PAP Diameter of Cells - Overview

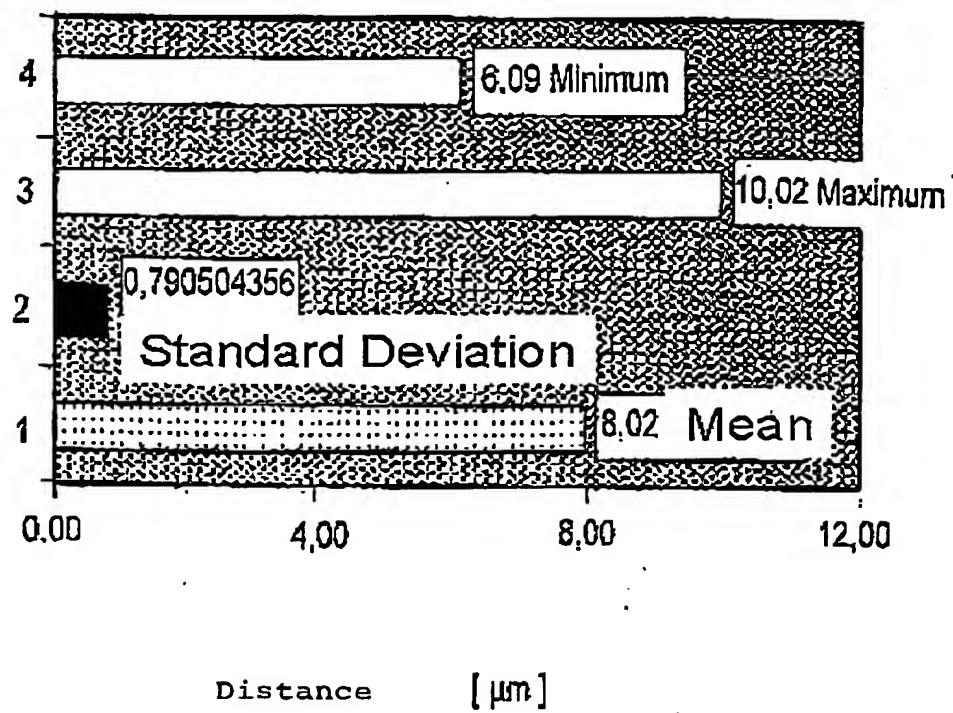


FIG.1A

## PAP Diameter of Cells - Overview

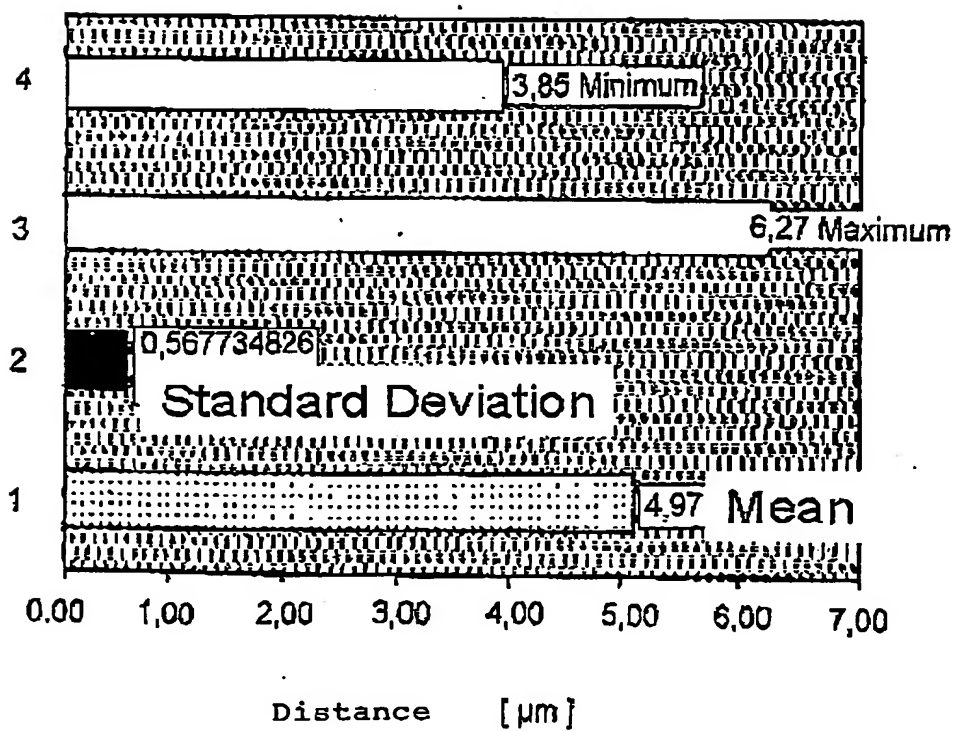
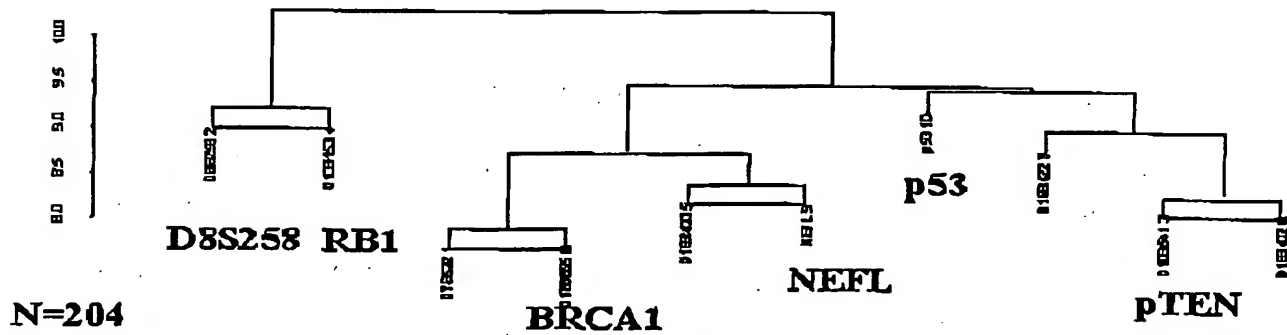
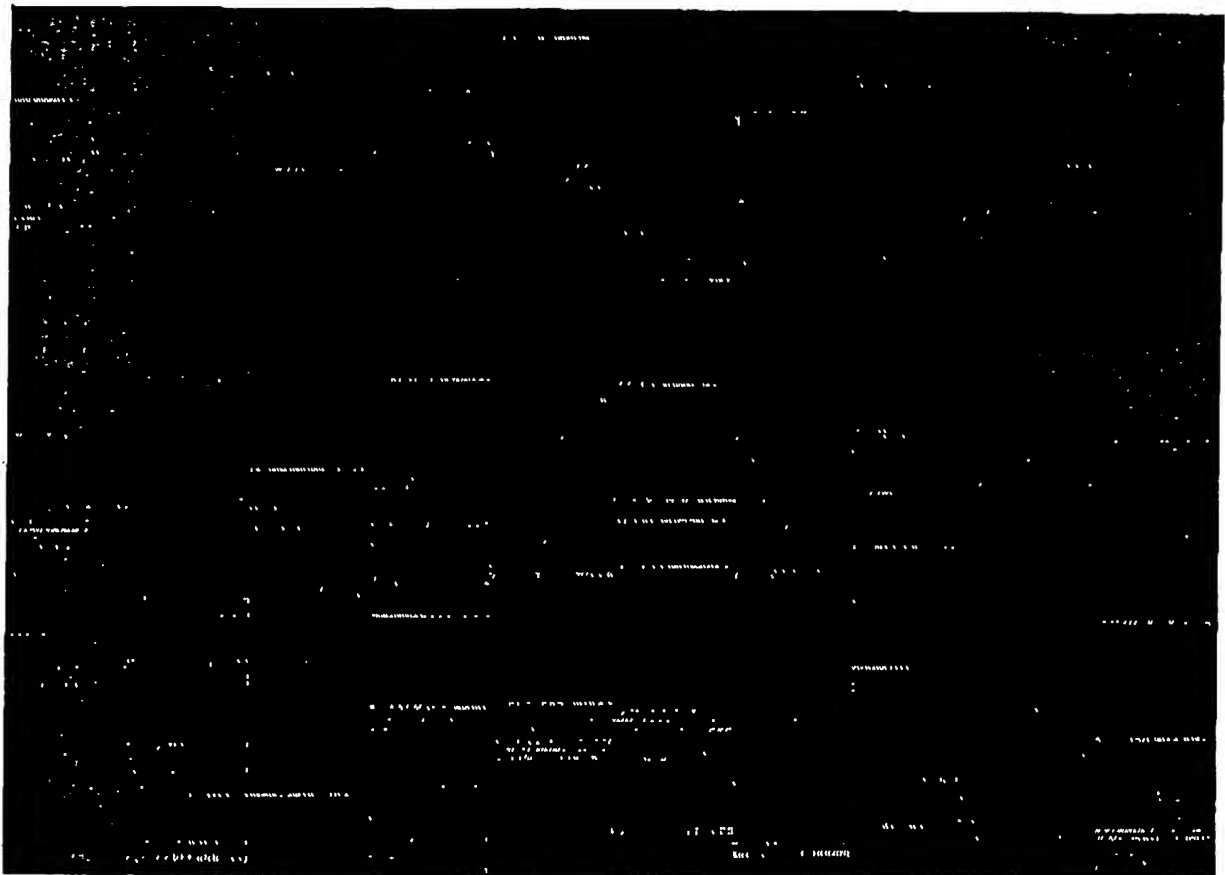


FIG.1B

# Cluster Analysis of Microsatellite Aberrations of Organ-Confined Primary Prostate Cancer



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Observations

# Multiple Genetic Progression of Prostate Cancer

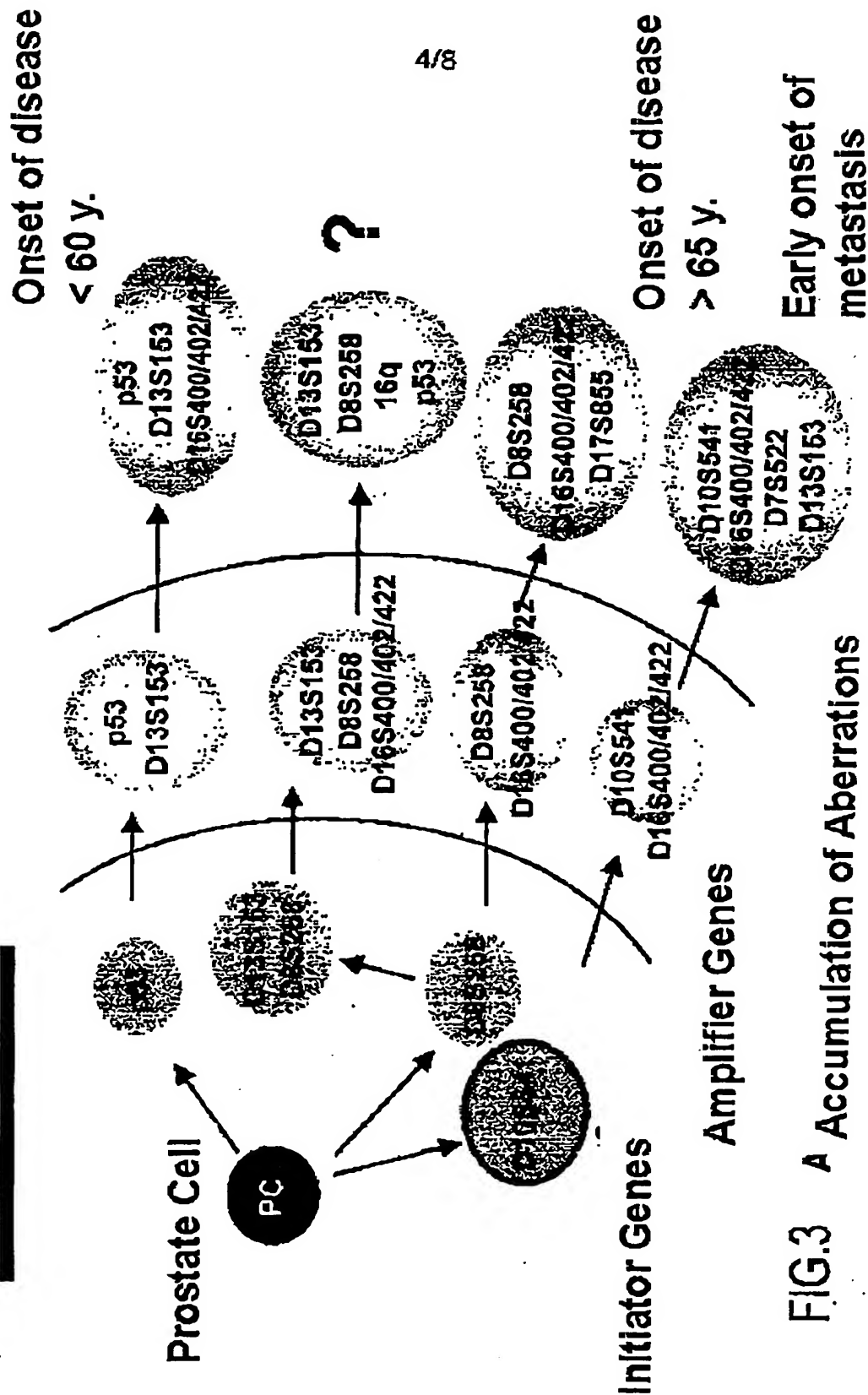


FIG.3 A Accumulation of Aberrations

Primary Tumor in the Prostate

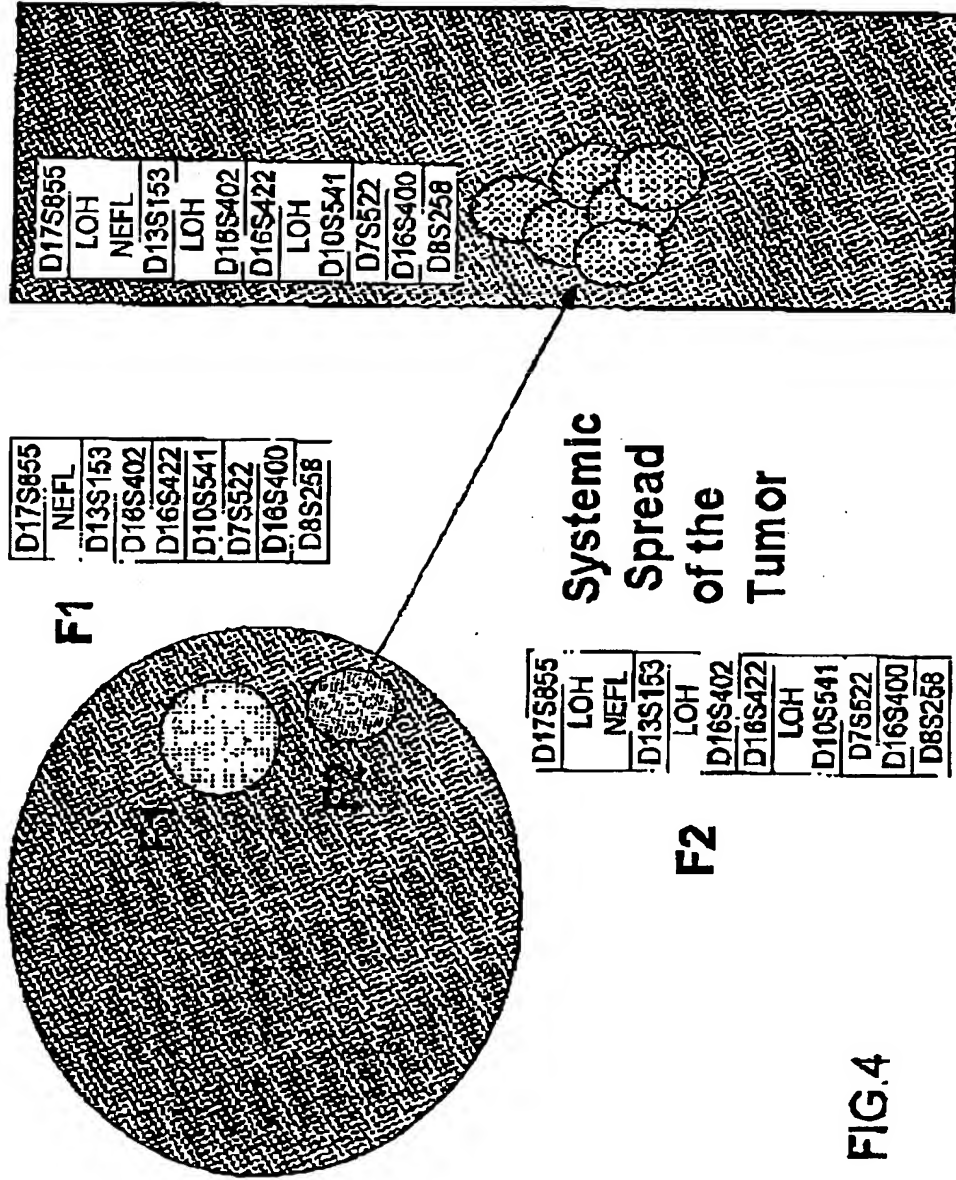


FIG. 4

# Genomic Aberrations of Circulating Cancer Cells

Two way hierarchical clustering  
 Data: RERLOHneuauswertungCluster[x.CC.]  
 Variables: 30CC 1 ... 117CC  
 Number of Variables: 24 Observables: 9  
 Model: hclust ( hierarchical clustering )  
 Method: compact ( complete linkage )  
 Distance Metric: euclidean  
 Data scaled: No

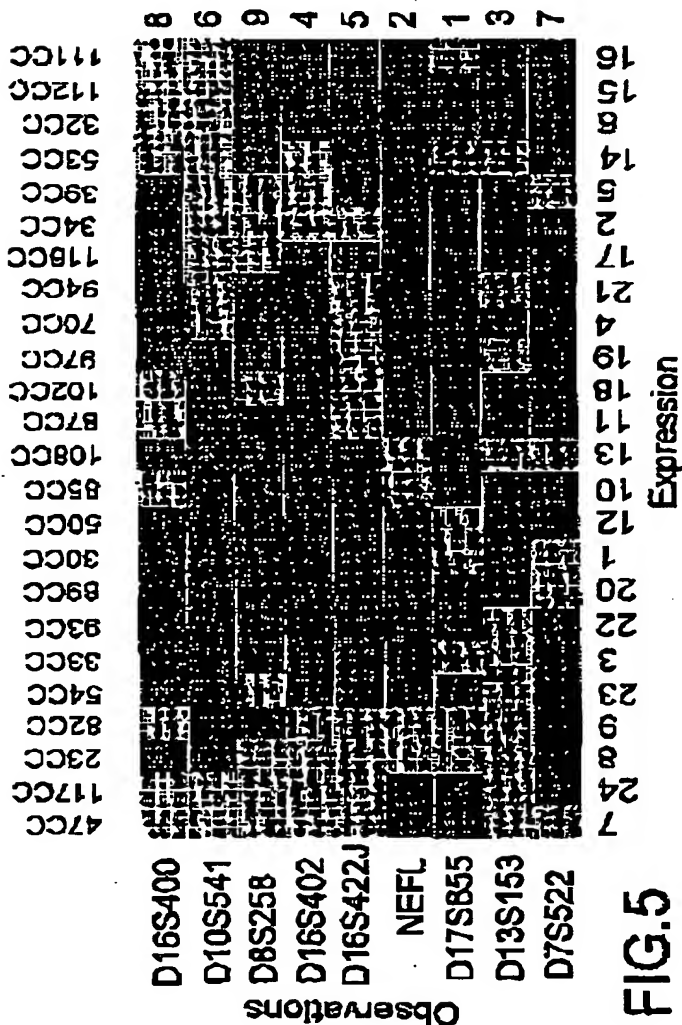
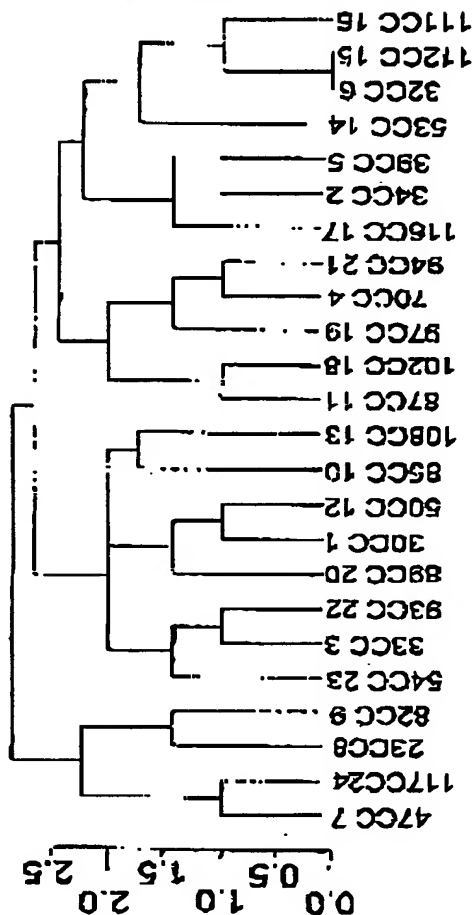


FIG.5

# Disease-free Survival of Prostate Cancer Patients and Aberrations at Chromosome 8p

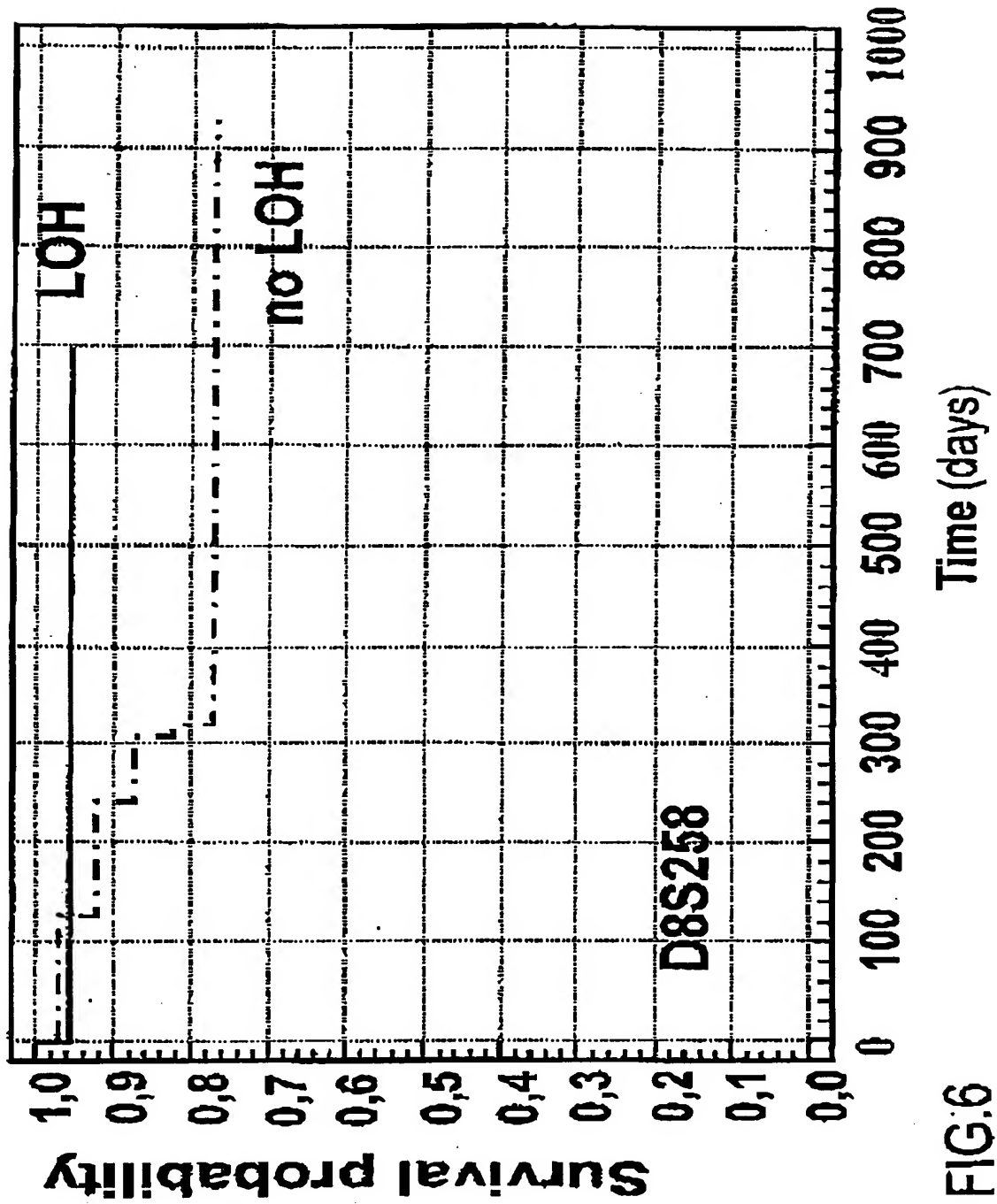


FIG:6

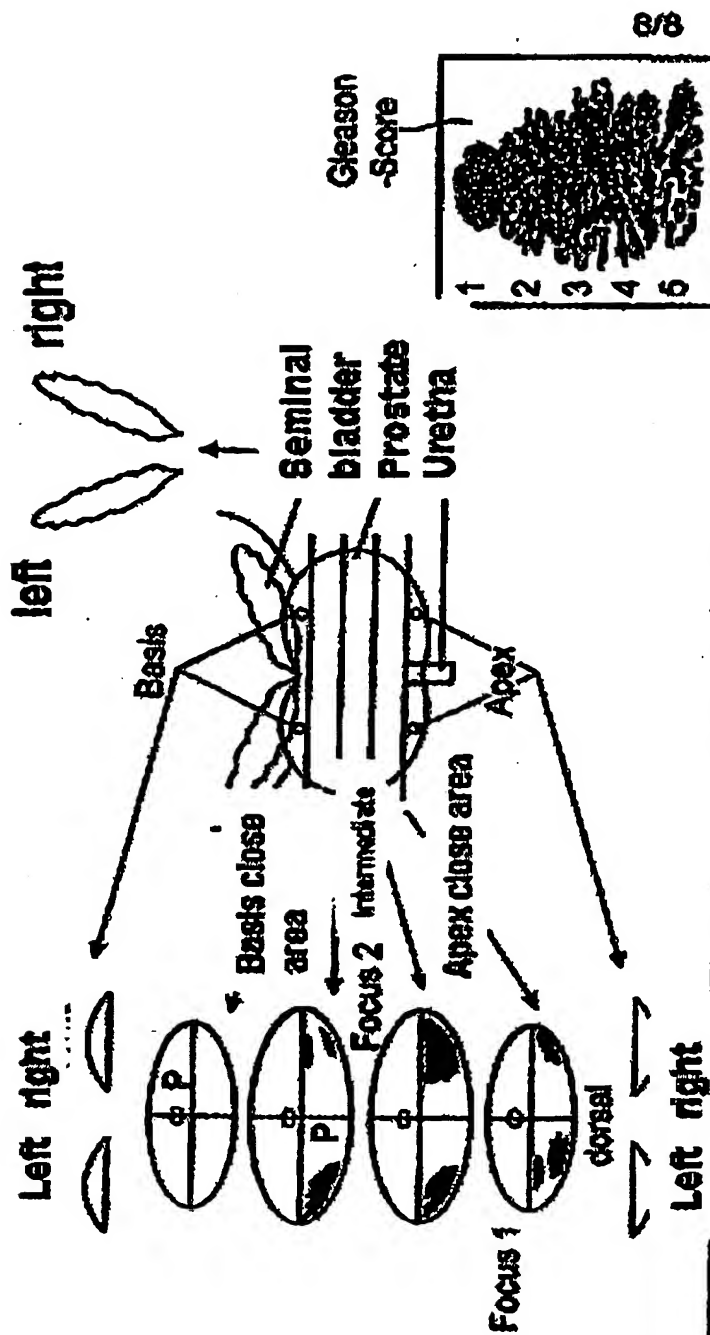


FIG. 7

<p><b>P</b> Prostatic intraepithelial Neoplasia (PIN)</p> <p> Adenocarcinoma</p> <p><b>—</b> Capsula invasion</p> <p><b>==</b> Extraprostatic spread</p> <p><b>*</b> Positive Margins</p>	<p><b>Pattern of Differentiation</b></p> <p>Highly differentiated 0</p> <p>Poor differentiated 1</p> <p>Cribriform 2</p> <p>Solid 3</p> <p>la = 0 lb = 1 lla = 2 llb = 3 llla = 4 llb = 5</p>	<p><b>Grade of Malignancy (Helpap)</b></p> <p>3 + 3 = 6</p> <p>lla</p>
<p>pT: 2b</p>	<p><b>Gleason Score</b></p>	<p><b>Grade of Malignancy (Helpap)</b></p>

Prostate Cancer Land Map



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